

- (d) backtranslating each of said (poly)peptide sequences into a corresponding coding nucleic acid sequence;
- (e) setting up cleavage sites in regions adjacent to or between the ends of sub-sequences encoding said sub-elements, each of said cleavage sites:
  - (ea) being unique within each of said coding nucleic acid sequences;
  - (eb) being common to the corresponding sub-sequences of any said coding nucleic acids.

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P1*
- 57. A collection of (poly)peptides comprising a plurality of (poly)peptides according to claim 56.
  - 58. A kit comprising a (poly)peptide according to claim 56.
  - 59. A kit comprising a collection of (poly)peptides according to claim 57.
  - 60. A collection of (poly)peptides according to claim 57, comprising specific (poly)peptides wherein the genes encoding said specific (poly)peptides
    - (a) are either homologous, or represent consensus gene sequences derived from at least three homologous genes, and
    - (b) carry cleavage sites, each of which:
      - (ba) lie at or adjacent to the ends of genetic sub-sequences which encode structural sub-elements,
      - (bb) are unique within each gene sequence,
      - (bc) do not form compatible sites with respect to any single sub-sequence, and
      - (bd) are common to all homologous sub-sequences.
  - 61. A kit comprising a collection of (poly)peptides according to claim 60.

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